

SEQUENCE LISTING

#11/4B

<110> Ashkenazi, Avi J.
Baker, Kevin
Gurney, Austin
Wood, William

<120> Apo-2DcR

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<140> US 08/878,168

<141> 1997-06-18

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<211> 259

<212> PRT

<213> HUMAN

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Glu	Val	Pro	Gln	Gln	Thr	Val	Ala	Pro	Gln	Gln	Gln	Arg	His	Ser	Phe
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Lys	Gly	Glu	Glu	Cys	Pro	Ala	Gly	Ser	His	Arg	Ser	Glu	His	Thr	Gly
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Ala	Cys	Asn	Pro	Cys	Thr	Glu	Gly	Val	Asp	Tyr	Thr	Asn	Ala	Ser	Asn
65					70				75					80	
Asn	Glu	Pro	Ser	Cys	Phe	Pro	Cys	Thr	Val	Cys	Lys	Ser	Asp	Gln	Lys
			85						90					95	
His	Lys	Ser	Ser	Cys	Thr	Met	Thr	Arg	Asp	Thr	Val	Cys	Gln	Cys	Lys
			100					105					110		
Glu	Gly	Thr	Phe	Arg	Asn	Glu	Asn	Ser	Pro	Glu	Met	Cys	Arg	Lys	Cys
		115					120					125			
Ser	Arg	Cys	Pro	Ser	Gly	Glu	Val	Gln	Val	Ser	Asn	Cys	Thr	Ser	Trp
	130				135						140				
Asp	Asp	Ile	Gln	Cys	Val	Glu	Glu	Phe	Gly	Ala	Asn	Ala	Thr	Val	Glu
145					150					155					160
Thr	Pro	Ala	Ala	Glu	Glu	Thr	Met	Asn	Thr	Ser	Pro	Gly	Thr	Pro	Ala
			165						170					175	
Pro	Ala	Ala	Glu	Glu	Thr	Met	Asn	Thr	Ser	Pro	Gly	Thr	Pro	Ala	Pro
			180					185					190		
Ala	Ala	Glu	Glu	Thr	Met	Thr	Thr	Ser	Pro	Gly	Thr	Pro	Ala	Pro	Ala
	195						200					205			
Ala	Glu	Glu	Thr	Met	Thr	Thr	Ser	Pro	Gly	Thr	Pro	Ala	Pro	Ala	Ala

210	Glu	Glu	Thr	Met	Thr	Thr	215	Ser	Pro	Gly	Thr	Pro	220	Ala	Ser	Ser	His	Tyr
225	Leu	Ser	Cys	Thr	Ile	Val	230	Gly	Ile	Ile	Val	Leu	235	Ile	Val	Leu	Leu	Ile
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agagcgcccc	ggccgcctga	tggccgaggc	aggggtgcgc	ccaggaccca	ggacggcgtc	180
gggaaccata	cc atg gcc	cgg atc ccc	aag acc cta	aag ttc gtc	gtc gtc gtc	231
	Met Ala Arg	Ile Pro Lys	Thr Leu Lys	Phe Val Val	Val Val Val	
	1		5		10	

atc gtc gcg gtc ctg ctg cca gtc cta gct tac tct gcc acc act gcc	279
Ile Val Ala Val Leu Leu Pro Val Leu Ala Tyr Ser Ala Thr Thr Ala	
15 20 25	

cgg cag gag gaa gtt ccc cag cag aca gtg gcc cca cag caa cag agg	327
Arg Gln Glu Glu Val Pro Gln Gln Thr Val Ala Pro Gln Gln Gln Arg	
30 35 40 45	

cac agc ttc aag ggg gag gag tgt cca gca gga tct cat aga tca gaa	375
His Ser Phe Lys Gly Glu Glu Cys Pro Ala Gly Ser His Arg Ser Glu	
50 55 60	

cat act gga gcc tgt aac ccg tgc aca gag ggt gtg gat tac acc aac	423
His Thr Gly Ala Cys Asn Pro Cys Thr Glu Gly Val Asp Tyr Thr Asn	
65 70 75	

gct tcc aac aat gaa cct tct tgc ttc cca tgt aca gtt tgt aaa tca	471
Ala Ser Asn Asn Glu Pro Ser Cys Phe Pro Cys Thr Val Cys Lys Ser	
80 85 90	

gat caa aaa cat aaa agt tcc tgc acc atg acc aga gac aca gtg tgt	519
Asp Gln Lys His Lys Ser Ser Cys Thr Met Thr Arg Asp Thr Val Cys	
95 100 105	

cag tgt aaa gaa ggc acc ttc cgg aat gaa aac tcc cca gag atg tgc	567
Gln Cys Lys Glu Gly Thr Phe Arg Asn Glu Asn Ser Pro Glu Met Cys	
110 115 120 125	

cgg aag tgt agc agg tgc cct agt ggg gaa gtc caa gtc agt aat tgt	615
Arg Lys Cys Ser Arg Cys Pro Ser Gly Glu Val Gln Val Ser Asn Cys	
130 135 140	
acg tcc tgg gat gat atc cag tgt gtt gaa gaa ttt ggt gcc aat gcc	663
Thr Ser Trp Asp Asp Ile Gln Cys Val Glu Glu Phe Gly Ala Asn Ala	
145 150 155	
act gtg gaa acc cca gct gct gaa gag aca atg aac acc agc ccg ggg	711
Thr Val Glu Thr Pro Ala Ala Glu Glu Thr Met Asn Thr Ser Pro Gly	
160 165 170	
act cct gcc cca gct gct gaa gag aca atg aac acc agc cca ggg act	759
Thr Pro Ala Pro Ala Ala Glu Glu Thr Met Asn Thr Ser Pro Gly Thr	
175 180 185	
cct gcc cca gct gct gaa gag aca atg acc acc agc ccg ggg act cct	807
Pro Ala Pro Ala Ala Glu Glu Thr Met Thr Thr Ser Pro Gly Thr Pro	
190 195 200 205	
gcc cca gct gct gaa gag aca atg acc acc agc ccg ggg act cct gcc	855
Ala Pro Ala Ala Glu Glu Thr Met Thr Thr Ser Pro Gly Thr Pro Ala	
210 215 220	
cca gct gct gaa gag aca atg acc acc agc ccg ggg act cct gcc tct	903
Pro Ala Ala Glu Glu Thr Met Thr Thr Ser Pro Gly Thr Pro Ala Ser	
225 230 235	
tct cat tac ctc tca tgc acc atc gta ggg atc ata gtt cta att gtg	951
Ser His Tyr Leu Ser Cys Thr Ile Val Gly Ile Ile Val Leu Ile Val	
240 245 250	
ctt ctg att gtg ttt gtt tgaaagactt cactgtggaa gaaattcctt	999
Leu Leu Ile Val Phe Val	
255	
ccttacctga aaggttcagg taggcgctgg ctgagggcgg ggggcgctgg acactctctg	1059
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aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa	1179
a	1180

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 20 25 30
 Gln Asp Gly Val Gly Asn His Thr Met Ala Arg Ile Pro Lys Thr Leu
 35 40 45

Lys	Phe	Val	Val	Val	Ile	Val	Ala	Val	Leu	Leu	Pro	Val	Leu	Ala	Tyr
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Ser	Ala	Thr	Thr	Ala	Arg	Gln	Glu	Glu	Val	Pro	Gln	Gln	Thr	Val	Ala
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Pro	Gln	Gln	Gln	Arg	His	Ser	Phe	Lys	Gly	Glu	Glu	Cys	Pro	Ala	Gly
				85					90					95	
Ser	His	Arg	Ser	Glu	His	Thr	Gly	Ala	Cys	Asn	Pro	Cys	Thr	Glu	Gly
			100					105					110		
Val	Asp	Tyr	Thr	Asn	Ala	Ser	Asn	Asn	Glu	Pro	Ser	Cys	Phe	Pro	Cys
	115						120					125			
Thr	Val	Cys	Lys	Ser	Asp	Gln	Lys	His	Lys	Ser	Ser	Cys	Thr	Met	Thr
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Arg	Asp	Thr	Val	Cys	Gln	Cys	Lys	Glu	Gly	Thr	Phe	Arg	Asn	Glu	Asn
145					150					155					160
Ser	Pro	Glu	Met	Cys	Arg	Lys	Cys	Ser	Arg	Cys	Pro	Ser	Gly	Glu	Val
				165					170					175	
Gln	Val	Ser	Asn	Cys	Thr	Ser	Trp	Asp	Asp	Ile	Gln	Cys	Val	Glu	Glu
			180					185					190		
Phe	Gly	Ala	Asn	Ala	Thr	Val	Glu	Thr	Pro	Ala	Ala	Glu	Glu	Thr	Met
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Asn	Thr	Ser	Pro	Gly	Thr	Pro	Ala	Pro	Ala	Ala	Glu	Glu	Thr	Met	Asn
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Thr	Ser	Pro	Gly	Thr	Pro	Ala	Pro	Ala	Ala	Glu	Glu	Thr	Met	Thr	Thr
225					230					235					240
Ser	Pro	Gly	Thr	Pro	Ala	Pro	Ala	Ala	Glu	Glu	Thr	Met	Thr	Thr	Ser
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Pro	Gly	Thr	Pro	Ala	Pro	Ala	Ala	Glu	Glu	Thr	Met	Thr	Thr	Ser	Pro
			260					265					270		
Gly	Thr	Pro	Ala	Ser	Ser	His	Tyr	Leu	Ser	Cys	Thr	Ile	Val	Gly	Ile
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<220>
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Met Gln Gly Val Lys Glu Arg Phe Leu Pro Leu Gly Asn															
-40					-35					-30					
tct ggg gac aga gcg ccc cgg ccg cct gat ggc cga ggc agg gtg cga															159
Ser Gly Asp Arg Ala Pro Arg Pro Pro Asp Gly Arg Gly Arg Val Arg															

-25					-20					-15						
ccc	agg	acc	cag	gac	ggc	gtc	ggg	aac	cat	acc	atg	gcc	cgg	atc	ccc	207
Pro	Arg	Thr	Gln	Asp	Gly	Val	Gly	Asn	His	Thr	Met	Ala	Arg	Ile	Pro	
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Lys	Thr	Leu	Lys	Phe	Val	Val	Val	Ile	Val	Ala	Val	Leu	Leu	Pro	Val	
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Leu	Ala	Tyr	Ser	Ala	Thr	Thr	Ala	Arg	Gln	Glu	Glu	Val	Pro	Gln	Gln	
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aca	gtg	gcc	cca	cag	caa	cag	agg	cac	agc	ttc	aag	ggg	gag	gag	tgt	351
Thr	Val	Ala	Pro	Gln	Gln	Gln	Arg	His	Ser	Phe	Lys	Gly	Glu	Glu	Cys	
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cca	gca	gga	tct	cat	aga	tca	gaa	cat	act	gga	gcc	tgt	aac	ccg	tgc	399
Pro	Ala	Gly	Ser	His	Arg	Ser	Glu	His	Thr	Gly	Ala	Cys	Asn	Pro	Cys	
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aca	gag	ggt	gtg	gat	tac	acc	aac	gct	tcc	aac	aat	gaa	cct	tct	tgc	447
Thr	Glu	Gly	Val	Asp	Tyr	Thr	Asn	Ala	Ser	Asn	Asn	Glu	Pro	Ser	Cys	
70					75					80					85	
ttc	cca	tgt	aca	gtt	tgt	aaa	tca	gat	caa	aaa	cat	aaa	agt	tcc	tgc	495
Phe	Pro	Cys	Thr	Val	Cys	Lys	Ser	Asp	Gln	Lys	His	Lys	Ser	Ser	Cys	
				90					95					100		
acc	atg	acc	aga	gac	aca	gtg	tgt	cag	tgt	aaa	gaa	ggc	acc	ttc	cgg	543
Thr	Met	Thr	Arg	Asp	Thr	Val	Cys	Gln	Cys	Lys	Glu	Gly	Thr	Phe	Arg	
			105					110					115			
aat	gaa	aac	tcc	cca	gag	atg	tgc	cgg	aag	tgt	agc	agg	tgc	cct	agt	591
Asn	Glu	Asn	Ser	Pro	Glu	Met	Cys	Arg	Lys	Cys	Ser	Arg	Cys	Pro	Ser	
		120					125					130				
ggg	gaa	gtc	caa	gtc	agt	aat	tgt	acg	tcc	tgg	gat	gat	atc	cag	tgt	639
Gly	Glu	Val	Gln	Val	Ser	Asn	Cys	Thr	Ser	Trp	Asp	Asp	Ile	Gln	Cys	
		135				140					145					
gtt	gaa	gaa	ttt	ggt	gcc	aat	gcc	act	gtg	gaa	acc	cca	gct	gct	gaa	687
Val	Glu	Glu	Phe	Gly	Ala	Asn	Ala	Thr	Val	Glu	Thr	Pro	Ala	Ala	Glu	
150					155					160					165	
gag	aca	atg	aac	acc	agc	ccg	ggg	act	cct	gcc	cca	gct	gct	gaa	gag	735
Glu	Thr	Met	Asn	Thr	Ser	Pro	Gly	Thr	Pro	Ala	Pro	Ala	Ala	Glu	Glu	
				170				175						180		
aca	atg	aac	acc	agc	cca	ggg	act	cct	gcc	cca	gct	gct	gaa	gag	aca	783
Thr	Met	Asn	Thr	Ser	Pro	Gly	Thr	Pro	Ala	Pro	Ala	Ala	Glu	Glu	Thr	
			185					190					195			

atg acc acc agc ccg ggg act cct gcc cca gct gct gaa gag aca atg	831
Met Thr Thr Ser Pro Gly Thr Pro Ala Pro Ala Ala Glu Glu Thr Met	
200 205 210	

acc acc agc ccg ggg act cct gcc cca gct gct gaa gag aca atg acc	879
Thr Thr Ser Pro Gly Thr Pro Ala Pro Ala Ala Glu Glu Thr Met Thr	
215 220 225	

acc agc ccg ggg act cct gcc tct tct cat tac ctc tca tgc acc atc	927
Thr Ser Pro Gly Thr Pro Ala Ser Ser His Tyr Leu Ser Cys Thr Ile	
230 235 240 245	

gta ggg atc ata gtt cta att gtg ctt ctg att gtg ttt gtt	969
Val Gly Ile Ile Val Leu Ile Val Leu Leu Ile Val Phe Val	
250 255	

tgaaagactt cactgtggaa gaaattcctt ccttacctga aaggttcagg taggcgctgg	1029
ctgagggcgcg ggggcgctgg acactctctg ccctgcctcc ctctgctgtg ttcccacaga	1089
cagaaacgcc tgcccctgcc ccaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa	1149
aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa a	1180

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35 40 45	
Lys	

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 Lys Ser Ser Cys Thr Met Thr Arg Asp Thr Val Cys Gln Cys Lys Glu
 35 40 45

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 aggcagcggg 70

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 aagagcgttc cctaccgcc atg gaa caa cgg gga cag aac gcc ccg gcc gct 172
 Met Glu Gln Arg Gly Gln Asn Ala Pro Ala Ala
 1 5 10

tcg ggg gcc cgg aaa agg cac ggc cca gga ccc agg gag gcg cgg gga 220
 Ser Gly Ala Arg Lys Arg His Gly Pro Gly Pro Arg Glu Ala Arg Gly
 15 20 25

gcc agg cct ggg ctc cgg gtc ccc aag acc ctt gtg ctc gtt gtc gcc 268
 Ala Arg Pro Gly Leu Arg Val Pro Lys Thr Leu Val Leu Val Val Ala
 30 35 40

gcg gtc ctg ctg ttg gtc tca gct gag tct gct ctg atc acc caa caa 316
 Ala Val Leu Leu Leu Val Ser Ala Glu Ser Ala Leu Ile Thr Gln Gln
 45 50 55

gac cta gct ccc cag cag aga gcg gcc cca caa caa aag agg tcc agc 364
 Asp Leu Ala Pro Gln Gln Arg Ala Ala Pro Gln Gln Lys Arg Ser Ser
 60 65 70 75

ccc	tca	gag	gga	ttg	tgt	cca	cct	gga	cac	cat	atc	tca	gaa	gac	ggt	412
Pro	Ser	Glu	Gly	Leu	Cys	Pro	Pro	Gly	His	His	Ile	Ser	Glu	Asp	Gly	
				80					85					90		
aga	gat	tgc	atc	tcc	tgc	aaa	tat	gga	cag	gac	tat	agc	act	cac	tgg	460
Arg	Asp	Cys	Ile	Ser	Cys	Lys	Tyr	Gly	Gln	Asp	Tyr	Ser	Thr	His	Trp	
			95					100					105			
aat	gac	ctc	ctt	ttc	tgc	ttg	cgc	tgc	acc	agg	tgt	gat	tca	ggt	gaa	508
Asn	Asp	Leu	Leu	Phe	Cys	Leu	Arg	Cys	Thr	Arg	Cys	Asp	Ser	Gly	Glu	
		110					115					120				
gtg	gag	cta	agt	ccc	tgc	acc	acg	acc	aga	aac	aca	gtg	tgt	cag	tgc	556
Val	Glu	Leu	Ser	Pro	Cys	Thr	Thr	Thr	Arg	Asn	Thr	Val	Cys	Gln	Cys	
	125					130					135					
gaa	gaa	ggc	acc	ttc	cgg	gaa	gaa	gat	tct	cct	gag	atg	tgc	cgg	aag	604
Glu	Glu	Gly	Thr	Phe	Arg	Glu	Glu	Asp	Ser	Pro	Glu	Met	Cys	Arg	Lys	
140					145					150					155	
tgc	cgc	aca	ggg	tgt	ccc	aga	ggg	atg	gtc	aag	gtc	ggt	gat	tgt	aca	652
Cys	Arg	Thr	Gly	Cys	Pro	Arg	Gly	Met	Val	Lys	Val	Gly	Asp	Cys	Thr	
				160					165					170		
ccc	tgg	agt	gac	atc	gaa	tgt	gtc	cac	aaa	gaa	tca	ggc	atc	atc	ata	700
Pro	Trp	Ser	Asp	Ile	Glu	Cys	Val	His	Lys	Glu	Ser	Gly	Ile	Ile	Ile	
			175					180					185			
gga	gtc	aca	gtt	gca	gcc	gta	gtc	ttg	att	gtg	gct	gtg	ttt	gtt	tgc	748
Gly	Val	Thr	Val	Ala	Ala	Val	Val	Leu	Ile	Val	Ala	Val	Phe	Val	Cys	
		190					195					200				
aag	tct	tta	ctg	tgg	aag	aaa	gtc	ctt	cct	tac	ctg	aaa	ggc	atc	tgc	796
Lys	Ser	Leu	Leu	Trp	Lys	Lys	Val	Leu	Pro	Tyr	Leu	Lys	Gly	Ile	Cys	
	205					210					215					
tca	ggt	ggt	ggt	ggg	gac	cct	gag	cgt	gtg	gac	aga	agc	tca	caa	cga	844
Ser	Gly	Gly	Gly	Gly	Asp	Pro	Glu	Arg	Val	Asp	Arg	Ser	Ser	Gln	Arg	
220					225					230					235	
cct	ggg	gct	gag	gac	aat	gtc	ctc	aat	gag	atc	gtg	agt	atc	ttg	cag	892
Pro	Gly	Ala	Glu	Asp	Asn	Val	Leu	Asn	Glu	Ile	Val	Ser	Ile	Leu	Gln	
				240					245					250		
ccc	acc	cag	gtc	cct	gag	cag	gaa	atg	gaa	gtc	cag	gag	cca	gca	gag	940
Pro	Thr	Gln	Val	Pro	Glu	Gln	Glu	Met	Glu	Val	Gln	Glu	Pro	Ala	Glu	
			255					260					265			
cca	aca	ggt	gtc	aac	atg	ttg	tcc	ccc	ggg	gag	tca	gag	cat	ctg	ctg	988
Pro	Thr	Gly	Val	Asn	Met	Leu	Ser	Pro	Gly	Glu	Ser	Glu	His	Leu	Leu	
		270					275					280				

gaa ccg gca gaa gct gaa agg tct cag agg agg agg ctg ctg gtt cca	1036
Glu Pro Ala Glu Ala Glu Arg Ser Gln Arg Arg Arg Leu Leu Val Pro	
285 290 295	
gca aat gaa ggt gat ccc act gag act ctg aga cag tgc ttc gat gac	1084
Ala Asn Glu Gly Asp Pro Thr Glu Thr Leu Arg Gln Cys Phe Asp Asp	
300 305 310 315	
ttt gca gac ttg gtg ccc ttt gac tcc tgg gag ccg ctc atg agg aag	1132
Phe Ala Asp Leu Val Pro Phe Asp Ser Trp Glu Pro Leu Met Arg Lys	
320 325 330	
ttg ggc ctc atg gac aat gag ata aag gtg gct aaa gct gag gca gcg	1180
Leu Gly Leu Met Asp Asn Glu Ile Lys Val Ala Lys Ala Glu Ala Ala	
335 340 345	
ggc cac agg gac acc ttg tac acg atg ctg ata aag tgg gtc aac aaa	1228
Gly His Arg Asp Thr Leu Tyr Thr Met Leu Ile Lys Trp Val Asn Lys	
350 355 360	
acc ggg cga gat gcc tct gtc cac acc ctg ctg gat gcc ttg gag acg	1276
Thr Gly Arg Asp Ala Ser Val His Thr Leu Leu Asp Ala Leu Glu Thr	
365 370 375	
ctg gga gag aga ctt gcc aag cag aag att gag gac cac ttg ttg agc	1324
Leu Gly Glu Arg Leu Ala Lys Gln Lys Ile Glu Asp His Leu Leu Ser	
380 385 390 395	
tct gga aag ttc atg tat cta gaa ggt aat gca gac tct gcc wtg tcc	1372
Ser Gly Lys Phe Met Tyr Leu Glu Gly Asn Ala Asp Ser Ala Xaa Ser	
400 405 410	
taagtgtgat tctcttcagg aagtgagacc ttccctgggt tacctttttt ctggaaaaag	1432
cccaactgga ctccagtcag taggaaagt ccacaattgt cacatgaccg gtactggaag	1492
aaactctccc atccaacatc acccagtgga tggaacatcc tgtaactttt cactgcactt	1552
ggcattatatt ttataagctg aatgtgataa taaggacact atggaaatgt ctggatcatt	1612
ccgtttgtgc gtactttgag atttggtttt ggatgtcatt gttttcacag cactttttta	1672
tcctaataatgta aatgctttat ttattttattt gggctacatt gtaagatcca tctacaaaaa	1732
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 20 25 30
 Arg Val Pro Lys Thr Leu Val Leu Val Val Ala Ala Val Leu Leu Leu
 35 40 45

Val	Ser	Ala	Glu	Ser	Ala	Leu	Ile	Thr	Gln	Gln	Asp	Leu	Ala	Pro	Gln
50						55					60				
Gln	Arg	Ala	Ala	Pro	Gln	Gln	Lys	Arg	Ser	Ser	Pro	Ser	Glu	Gly	Leu
65					70					75					80
Cys	Pro	Pro	Gly	His	His	Ile	Ser	Glu	Asp	Gly	Arg	Asp	Cys	Ile	Ser
				85					90					95	
Cys	Lys	Tyr	Gly	Gln	Asp	Tyr	Ser	Thr	His	Trp	Asn	Asp	Leu	Leu	Phe
			100					105					110		
Cys	Leu	Arg	Cys	Thr	Arg	Cys	Asp	Ser	Gly	Glu	Val	Glu	Leu	Ser	Pro
		115					120					125			
Cys	Thr	Thr	Thr	Arg	Asn	Thr	Val	Cys	Gln	Cys	Glu	Glu	Gly	Thr	Phe
	130					135					140				
Arg	Glu	Glu	Asp	Ser	Pro	Glu	Met	Cys	Arg	Lys	Cys	Arg	Thr	Gly	Cys
145					150					155					160
Pro	Arg	Gly	Met	Val	Lys	Val	Gly	Asp	Cys	Thr	Pro	Trp	Ser	Asp	Ile
				165					170					175	
Glu	Cys	Val	His	Lys	Glu	Ser	Gly	Ile	Ile	Ile	Gly	Val	Thr	Val	Ala
			180					185					190		
Ala	Val	Val	Leu	Ile	Val	Ala	Val	Phe	Val	Cys	Lys	Ser	Leu	Leu	Trp
		195					200					205			
Lys	Lys	Val	Leu	Pro	Tyr	Leu	Lys	Gly	Ile	Cys	Ser	Gly	Gly	Gly	Gly
	210					215					220				
Asp	Pro	Glu	Arg	Val	Asp	Arg	Ser	Ser	Gln	Arg	Pro	Gly	Ala	Glu	Asp
225					230					235					240
Asn	Val	Leu	Asn	Glu	Ile	Val	Ser	Ile	Leu	Gln	Pro	Thr	Gln	Val	Pro
				245					250					255	
Glu	Gln	Glu	Met	Glu	Val	Gln	Glu	Pro	Ala	Glu	Pro	Thr	Gly	Val	Asn
			260					265					270		
Met	Leu	Ser	Pro	Gly	Glu	Ser	Glu	His	Leu	Leu	Glu	Pro	Ala	Glu	Ala
		275					280					285			
Glu	Arg	Ser	Gln	Arg	Arg	Arg	Leu	Leu	Val	Pro	Ala	Asn	Glu	Gly	Asp
	290					295					300				
Pro	Thr	Glu	Thr	Leu	Arg	Gln	Cys	Phe	Asp	Asp	Phe	Ala	Asp	Leu	Val
305					310					315					320
Pro	Phe	Asp	Ser	Trp	Glu	Pro	Leu	Met	Arg	Lys	Leu	Gly	Leu	Met	Asp
				325					330					335	
Asn	Glu	Ile	Lys	Val	Ala	Lys	Ala	Glu	Ala	Ala	Gly	His	Arg	Asp	Thr
			340					345					350		
Leu	Tyr	Thr	Met	Leu	Ile	Lys	Trp	Val	Asn	Lys	Thr	Gly	Arg	Asp	Ala
		355					360					365			
Ser	Val	His	Thr	Leu	Leu	Asp	Ala	Leu	Glu	Thr	Leu	Gly	Glu	Arg	Leu
	370					375					380				
Ala	Lys	Gln	Lys	Ile	Glu	Asp	His	Leu	Leu	Ser	Ser	Gly	Lys	Phe	Met
385					390					395					400
Tyr	Leu	Glu	Gly	Asn	Ala	Asp	Ser	Ala	Xaa	Ser					
				405					410						

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 <213> HUMAN

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29

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 aggatgggaa gtgtgtgata tacccttgat

30

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Arg	Ala	Arg	Ala	Gly	Arg	Ala	Pro	Gly	Pro	Pro	Pro	Ala	Arg	Glu	Ala
			20					25					30		
Ser	Pro	Arg	Leu	Arg	Val	His	Lys	Thr	Phe	Lys	Phe	Val	Val	Val	Gly
		35					40					45			
Val	Leu	Leu	Gln	Val	Val	Pro	Ser	Ser	Ala	Ala	Thr	Ile	Lys	Leu	His
	50				55						60				
Asp	Gln	Ser	Ile	Gly	Thr	Gln	Gln	Trp	Glu	His	Ser	Pro	Leu	Gly	Glu
65					70					75					80
Leu	Cys	Pro	Pro	Gly	Ser	His	Arg	Ser	Glu	Arg	Pro	Gly	Ala	Cys	Asn
				85					90					95	
Arg	Cys	Thr	Glu	Gly	Val	Gly	Tyr	Thr	Asn	Ala	Ser	Asn	Asn	Leu	Phe
			100					105					110		
Ala	Cys	Leu	Pro	Cys	Thr	Ala	Cys	Lys	Ser	Asp	Glu	Glu	Glu	Arg	Ser
		115					120					125			
Pro	Cys	Thr	Thr	Thr	Arg	Asn	Thr	Ala	Cys	Gln	Cys	Lys	Pro	Gly	Thr
	130					135					140				
Phe	Arg	Asn	Asp	Asn	Ser	Ala	Glu	Met	Cys	Arg	Lys	Cys	Ser	Thr	Gly
145					150					155					160
Cys	Pro	Arg	Gly	Met	Val	Lys	Val	Lys	Asp	Cys	Thr	Pro	Trp	Ser	Asp
				165					170					175	
Ile	Glu	Cys	Val	His	Lys	Glu	Ser	Gly	Asn	Gly	His	Asn	Ile	Trp	Val
			180					185					190		
Ile	Leu	Val	Val	Thr	Leu	Val	Val	Pro	Leu	Leu	Leu	Val	Ala	Val	Leu
		195					200					205			
Ile	Val	Cys	Cys	Cys	Ile	Gly	Ser	Gly	Cys	Gly	Gly	Asp	Pro	Lys	Cys
	210					215					220				
Met	Asp	Arg	Val	Cys	Phe	Trp	Arg	Leu	Gly	Leu	Leu	Arg	Gly	Pro	Gly
225					230					235					240
Ala	Glu	Asp	Asn	Ala	His	Asn	Glu	Ile	Leu	Ser	Asn	Ala	Asp	Ser	Leu
				245					250					255	
Ser	Thr	Phe	Val	Ser	Glu	Gln	Gln	Met	Glu	Ser	Gln	Glu	Pro	Ala	Asp
			260					265					270		
Leu	Thr	Gly	Val	Thr	Val	Gln	Ser	Pro	Gly	Glu	Ala	Gln	Cys	Leu	Leu
		275				280						285			
Gly	Pro	Ala	Glu	Ala	Glu	Gly	Ser	Gln	Arg	Arg	Arg	Leu	Leu	Val	Pro

290	Ala Asn Gly Ala Asp	295	Pro Thr Glu Thr Leu Met	300	Leu Phe Phe Asp Lys
305	Phe Ala Asn Ile Val	310	Pro Phe Asp Ser Trp	315	Asp Gln Leu Met Arg Gln
		325		330	
	Leu Asp Leu Thr Lys	335	Asn Glu Ile Asp Val	340	Val Arg Ala Gly Thr Ala
		345		350	
	Gly Pro Gly Asp Ala	360	Leu Tyr Ala Met Leu	365	Met Lys Trp Val Asn Lys
		370		375	
	Thr Gly Arg Asn Ala	380	Ser Ile His Thr Leu	385	Leu Asp Ala Leu Glu Arg
		390		395	
	Met Glu Glu Arg His	405	Ala Lys Glu Lys Ile	410	Gln Asp Leu Leu Val Asp
					415
	Ser Gly Lys Phe Ile		Tyr Leu Glu Asp Gly		Thr Gly Ser Ala Val Ser
	Leu Glu				

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	Leu Gly Leu Arg Glu Ala Glu Ile Glu Ala Val Glu Val Glu Ile Gly
	20 25 30
	Arg Phe Arg Asp Gln Gln Tyr Glu Met Leu Lys Arg Trp Arg Gln Gln
	35 40 45
	Gln Pro Ala Gly Leu Gly Ala Val Tyr Ala Ala Leu Glu Arg Met Gly
	50 55 60
	Leu Asp Gly Cys Val Glu Asp Leu Arg Ser
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	20 25 30
	Arg Cys Leu Arg Glu Ala Gln Tyr Ser Met Leu Ala Thr Trp Arg Arg
	35 40 45
	Arg Thr Pro Arg Arg Glu Ala Thr Leu Glu Leu Leu Gly Arg Val Leu
	50 55 60
	Arg Asp Met Asp Leu Leu Gly Cys Leu Glu Asp Ile Glu Glu
65	70 75

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<400> 17

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Asn	Gly	Val	Asn	Glu	Ala	Lys	Ile	Asp	Glu	Ile	Lys	Asn	Asp	Asn	Val
			20					25					30		
Gln	Asp	Thr	Ala	Glu	Gln	Lys	Val	Gln	Leu	Leu	Arg	Asn	Trp	His	Gln
		35					40					45			
Leu	His	Gly	Lys	Lys	Glu	Ala	Tyr	Asp	Thr	Leu	Ile	Lys	Asp	Leu	Lys
	50					55					60				
Lys	Ala	Asn	Leu	Cys	Thr	Leu	Ala	Glu	Lys	Ile	Gln	Thr			
65					70					75					